

SEQUENCE LISTING

<110> HABERMANN, PAUL
BENDER, RUDOLF

<120> SIGNAL SEQUENCES FOR PREPARING LEU-HIRUDIN BY SECRETION
BY E. COLI INTO THE CULTURE MEDIUM

<130> 02481.1693

<140>
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<160> 33

<170> PatentIn Ver. 2.1

<210> 1
<211> 25
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Oligonucleotide

<400> 1
tttttttaag cttgggctgc aggtc

25

<210> 2
<211> 54
<212> DNA
<213> Artificial Sequence

<220>
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tggcactggc aggttcgct accgtagcgc aagccttac gtatactgac tgca

54

<210> 3
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

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ttttttgaat tcatgaaaaa gacagctatc gcatttagcag tggcactggc aggttc

57

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<212> DNA
<213> Artificial Sequence

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<223> Description of Artificial Sequence: Primer

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<211> 56
<212> DNA
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<400> 5
ttttttgaat tcatgaaaaa cacctgggc ttggccattg gttctttat tgccgc 56

<210> 6
<211> 61
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 6
gttgccgtcg cagcgggcgt aatgtctgct caggcaatgg ctcttacgta tactgactgc 60
a 61

<210> 7
<211> 59
<212> DNA
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<400> 7
ttttttgaat tcatgatgat tactctgcgc aaacttcctc tggcggttgc cgtcgcagc 59

<210> 8
<211> 63
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer

<400> 8
ctaccctgat ggttaccgct ggtctgatgg gtaccgctgt tgctcttacg tataactgact 60
gca 63

<210> 9

<211> 60
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tttttgaat tcatgaaaaa aatgaacctg gctgtttgca tcgctaccct gatgggtacc 60

<210> 10
<211> 61
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer

<400> 10
ctgatcccg tctttgcagc gttctgcctg ccggtttcg cgcttacgta tactgactgc 60
a 61

<210> 11
<211> 56
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 11
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<210> 12
<211> 53
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 12
gctcccgctg ctgttcaccc cggttaccaa agcgcttacg tatactgact gca 53

<210> 13
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 13
tttttgaat tcatgaaaca gtcgaccatc gcgctggcgc tgctgccgt gctgttc 57

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<210> 14
<211> 53
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<220>
<223> Description of Artificial Sequence: Primer

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<400> 15
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<211> 64
<212> DNA
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tgca      64

<210> 17
<211> 65
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<400> 17
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atggc      65

<210> 18
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<212> DNA
<213> Artificial Sequence

<220>
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<223> Description of Artificial Sequence: Primer

<400> 18
tggtttcagc tttagtaaagc ggggttgcat ttgctttac gtataactgac tgcac 55

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<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 19
ttttggaaat tcataaaaaa gacaattatg tctctggctg tggtttcagc tttagtaagc 60

<210> 20
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence: Primer

<400> 20
cggcgctgag ttcgcctta ttttctcacc tatctttgc cttacgtat actgactgca 60

<210> 21
<211> 57
<212> DNA
<213> Artificial Sequence

<220>
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<400> 21
ttttttgaat tcatgtcatt tcatcaccgg gtatttaaac tgtcggcgct gagtctc 57

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<211> 227
<212> DNA
<213> Unknown Organism

<220>
<223> Description of Unknown Organism: Hirudin-encoding
DNA sequence

<220>
<221> CDS
<222> (1) .. (195)

<400> 22
 ctt acg tat act gac tgc act gaa tct ggt cag aac ctg tgc ctg tgc 48
 Leu Thr Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Leu Cys Leu Cys
 1 5 10 15

 gaa gga tct aac gtt tgc ggc cag ggt aac aaa tgc atc ctt gga tcc 96
 Glu Gly Ser Asn Val Cys Gly Gln Gly Asn Lys Cys Ile Leu Gly Ser
 20 25 30

 gac ggt gaa aag aac cag tgc gtt act ggc gaa ggt acc ccg aaa ccg 144
 Asp Gly Glu Lys Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Lys Pro
 35 40 45

 cag tct cat aac gac ggc gac ttc gaa gag atc cct gag gaa tac ctt 192
 Gln Ser His Asn Asp Gly Asp Phe Glu Glu Ile Pro Glu Glu Tyr Leu
 50 55 60

 cag taatagagct cgtcgacctg cagcccaagc tt 227
 Gln
 65

<210> 23
 <211> 65
 <212> PRT
 <213> Unknown Organism

 <220>
 <223> Description of Unknown Organism: Hirudin-encoded
 amino acid sequence

 <400> 23
 Leu Thr Tyr Thr Asp Cys Thr Glu Ser Gly Gln Asn Leu Cys Leu Cys 48
 1 5 10 15

 Glu Gly Ser Asn Val Cys Gly Gln Gly Asn Lys Cys Ile Leu Gly Ser
 20 25 30

 Asp Gly Glu Lys Asn Gln Cys Val Thr Gly Glu Gly Thr Pro Lys Pro
 35 40 45

 Gln Ser His Asn Asp Gly Asp Phe Glu Glu Ile Pro Glu Glu Tyr Leu
 50 55 60

 Gln
 65

<210> 24
 <211> 30
 <212> PRT
 <213> Unknown Organism

 <220>
 <223> Description of Unknown Organism: Control:
 cgtase-Ala-hirudin

<400> 24
Met Lys Arg Asn Arg Phe Phe Asn Thr Ser Ala Ala Ile Ala Ile Ser
1 5 10 15

Ile Ala Leu Asn Thr Phe Phe Cys Ser Met Gln Thr Ile Ala
20 25 30

<210> 25
<211> 21
<212> PRT
<213> Serratia marcescens

<220>
<223> Outer membrane protein

<400> 25
Met Lys Lys Thr Ala Ile Ala Leu Ala Val Ala Leu Ala Gly Phe Ala
1 5 10 15

Thr Val Ala Gln Ala
20

<210> 26
<211> 22
<212> PRT
<213> Pseudomonas fluorescens

<220>
<223> oprF protein

<400> 26
Met Lys Asn Thr Leu Gly Leu Ala Ile Gly Ser Leu Ile Ala Ala Thr
1 5 10 15

Ser Phe Gly Val Leu Ala
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<210> 27
<211> 25
<212> PRT
<213> Escherichia coli

<220>
<223> lamB protein

<400> 27
Met Met Ile Thr Leu Arg Lys Leu Pro Leu Ala Val Ala Val Ala Ala
1 5 10 15

Gly Val Met Ser Ala Gln Ala Met Ala
20 25

<210> 28
<211> 25

<212> PRT
<213> Shewanella putrefaciens

<220>
<223> Fumarate reductase

<400> 28
Met Lys Lys Met Asn Leu Ala Val Cys Ile Ala Thr Leu Met Gly Thr
1 5 10 15

Ala Gly Leu Met Gly Thr Ala Val Ala
20 25

<210> 29
<211> 23
<212> PRT
<213> Unknown Organism

<220>
<223> Description of Unknown Organism: Beta -
Lactamase/pBR322

<400> 29
Met Ser Ile Gln His Phe Arg Val Ala Leu Ile Pro Phe Phe Ala Ala
1 5 10 15

Phe Ser Leu Pro Val Phe Ala
20

<210> 30
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<212> PRT
<213> Escherichia coli

<220>
<223> Alk. phosphatase

<400> 30
Met Lys Gln Ser Thr Ile Ala Leu Ala Leu Leu Pro Leu Leu Phe Thr
1 5 10 15

Pro Val Thr Lys Ala
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<210> 31
<211> 21
<212> PRT
<213> Escherichia fergusonii

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<223> Alk. phosphatase

<400> 31
Met Lys Gln Ser Ala Ile Ala Leu Ala Leu Leu Ser Cys Leu Ile Thr
1 5 10 15

Pro Val Ser Gln Ala
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<210> 32
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<213> Paenibacillus macerans

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<223> Cyclodextrin glucanotransferase

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Met Lys Ser Arg Tyr Lys Arg Leu Thr Ser Leu Ala Leu Ser Leu Ser
1 5 10 15

Met Ala Leu Gly Ile Ser Leu Pro Ala Trp Ala
20 25

<210> 33
<211> 24
<212> PRT
<213> Salmonella typhimurium

<220>
<223> Outer membrane protein

<400> 33
Met Ser Phe His His Arg Val Phe Lys Leu Ser Ala Leu Ser Leu Ala
1 5 10 15

Leu Phe Ser His Leu Ser Phe Ala
20